(1/1)

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CYCD4     1                        MAEEN ELS LLC ESNVD EG IV..DE..TPIEI SI PQ GFS   39
CYCD2     1 MAENLACGETSESWIIDNDDDD NYGGGF NEIDYN HQ FAK DN..FGGNGSI PM GSS              58
CYCD1     1                 MRSYRFSDYLHM VS FSN MD FCGE SGVFSGES VDFSSSE              43
CYCD3     1 MAIRKEEESREEQSNSF LDA YCEEEKWD EGEEVEEN SSLSSSSSPFVV QQD                   55

CYCD4    40 QSE.SEEI IMEMV EKEKQH P DDYIKRLRSGDLD NVGRR ALNWIWKACEVHQ GP C             98
CYCD2    59 SSSLSE DRIKEM VREIEFCPGTDYVKRLLSGDLDLSV.RNQALDWILKVCAH HF GHLC            117
CYCD1    44 VDSWPGDS ACF EDERTFVPGHDYLSRFQTRS LDAS.AREDS AWILKVQAYYN QPLS            102
CYCD3    56 LFWEDEDL TLFSKE EQG SCLD.......DV LSTDRKEAVGWILRVNAHYG STLA               107

CYCD4    99 FC LAMNYLDRFL SVHD PSGKGW I LQLLAVACLSLAAK EETEVPML DLQVGDPQ FVFE           158
CYCD2   118 IC SMNYLDRFL TSYE PKD DWAA QLLAVACLSLASK EETDVPH   LQVE PKFVFE             177
CYCD1   103 AYLA NYMDRFL YARR PETSGWPMQLLAVACLSLAAK EEILVPS F FQVAGVKYLFE             162
CYCD3   108 AVLA I YLDKFI CSYS QRDKPWMLQLVSVACLSLAAK EETQVPL LLDFQVEE KYVFE            167

CYCD4   159 AKSVQRMELLVL NK KWRLRA TPCSYIR FLRKMSKCDQEPSNTL ISRSLQV AS TT KG           218
CYCD2   178 AKTIKRMELLVV T LNWRLQALTPFSFIDYFVDKIS...GHVSENL YRSSRFILNTT A               234
CYCD1   163 AKTIKRMELLVLS VLDWRLRS TPFDFISFFAYKIDL..RVPFSGS SPMLQSILSNIKE              220
CYCD3   168 AKTIQRMELLILS TLEWKMHL TP SFVDHI RRLGLKNNAHWD.F NKCHRL LSVISD             226

CYCD4   219 ID FLE RPSEAAAA AL SVSGE...LQRVHFDNS SFSPLFSL QKERVKK IGEMIES.DG            274
CYCD2   235 IEFLDFRPSE AAAAA SVSIS...GETECIDEEKALSSLIY KQERVKR LNLMRS TG               291
CYCD1   221 ASFLEYWPSS AAAA IL VANELPSLSSVVNPHES PETWCDG SKEKI VRCYRLMKAMAI            280
CYCD3   227 SR GYLPSV VAAATMMR I EQVDPFDP..LSYQ NLLGVLN TKEKVK CYDL LQ PV               284

CYCD4   275 SDLCS QT PNGVLEVSACCFSFK HD.S SSY HLS                                308
CYCD2   292 ENVRG SLSQEQA VAVRAVPASPVGVLEA CLSYRSEERTVESCTNSS QSSPDNNNNN   351
CYCD1   281 ENNRL T PKVIAKLRVSVRASS LTRPS DES SPCKRRKLSGYSWVGDE STSN        334
CYCD3   285 RICL QIQIQSSKKRKSHDSSS LNSP CVIDANPFNSDESSNDSWSASS CNPPTSSSSP   344

CYCD2   352 NNSKQEEEKTMRENKRVIHCLLQPKTTSTHDI 383
CYCD3   345 QQQPPLKKMRGAEENEKKKPILHLPWRIVATP 376
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Figure 1